RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/532, 0.28Source: PcTDate Processed by STIC: 04/11/2006

ENTERED

PCT

DATE: 04/11/2006

TIME: 12:37:36

```
Input Set : A:\65959-50.txt
                     Output Set: N:\CRF4\04112006\J532028.raw
      3 <110> APPLICANT: Boender, Piet
      5 <120> TITLE OF INVENTION: IMPROVED METHODS FOR GENERATING MULTIPLE RNA COPIES
     7 <130> FILE REFERENCE: 65959/50
      9 <140> CURRENT APPLICATION NUMBER: 10/532,028
     10 <141> CURRENT FILING DATE: 2005-04-21
     12 <150> PRIOR APPLICATION NUMBER: PCT/EP03/12058
    13 <151> PRIOR FILING DATE: 2003-10-30
    15 <150> PRIOR APPLICATION NUMBER: US 60/440,688
    16 <151> PRIOR FILING DATE: 2003-01-17
                                                                                    132
    18 <150> PRIOR APPLICATION NUMBER: EP 02447204.5
    19 <151> PRIOR FILING DATE: 2002-10-30
    21 <160> NUMBER OF SEQ ID NOS: 3
    23 <170> SOFTWARE: PatentIn version 3.3
    25 <210> SEQ ID NO: 1
    26 <211> LENGTH: 55
    27 <212> TYPE: DNA
    28 <213> ORGANISM: Artificial Sequence
    30 <220> FEATURE:
    31 <223> OTHER INFORMATION: Primer
    34 <220> FEATURE:
    35 <221> NAME/KEY: misc_feature
    36 <222> LOCATION: (50)..(55)
    37 <223> OTHER INFORMATION: N = an equimolar mixture of A, C, G and T bases of which the
last
     38
              nucleotide is a 2'-3' dideoxynucleotide and the final phosphate
             bond has been exchanged with a phosphorothicate bond
     41 <400> SEQUENCE: 1
W--> 42 aattetaata egaeteaeta tagggagaga aggataceae tagetagegn nnnnn
     45 <210> SEQ ID NO: 2
     46 <211> LENGTH: 36
     47 <212> TYPE: DNA
    48 <213> ORGANISM: Artificial Sequence
    50 <220> FEATURE:
    51 <223> OTHER INFORMATION: Primer
    54 <220> FEATURE:
    55 <221> NAME/KEY: misc feature
    56 <222> LOCATION: (31)..(36)
    57 <223> OTHER INFORMATION: N stands for an equimolar mixture of A, C, G and T bases - of
              which the last nucleotide is a 2'-3' dideoxynucleotide
    58
     60 <400> SEQUENCE: 2
W--> 61 aattctaata cgactcacta tagggagaga nnnnnn
                                                                               36
    64 <210> SEO ID NO: 3
    65 <211> LENGTH: 30
     66 <212> TYPE: DNA
```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/532,028

RAW SEQUENCE LISTING DATE: 04/11/2006
PATENT APPLICATION: US/10/532,028 TIME: 12:37:36

Input Set : A:\65959-50.txt

Output Set: N:\CRF4\04112006\J532028.raw

and the second of the second o

67 <213> ORGANISM: Artificial Sequence

69 <220> FEATURE:

70 <223> OTHER INFORMATION: Primer

72 <400> SEQUENCE: 3

73 agcacggaaa gacgatgacg gaaaaagaga

30

Contract of the second of the second

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 04/11/2006
PATENT APPLICATION: US/10/532,028 TIME: 12:37:37

Input Set : A:\65959-50.txt

Output Set: N:\CRF4\04112006\J532028.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 50,51,52,53,54,55
Seq#:2; N Pos. 31,32,33,34,35,36

VERIFICATION SUMMARY

. . . .

DATE: 04/11/2006

PATENT APPLICATION: US/10/532,028

TIME: 12:37:37

Input Set : A:\65959-50.txt

Output Set: N:\CRF4\04112006\J532028.raw

 $L\!:\!42~M\!:\!341~W\!:$ (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0 L:61 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0